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Range: from to Features: CDD

1: 11360168. Reports ...[gi:11360168] This record has been discontinued.

Comment Features Sequence

LOCUS T46372 859 aa linear PRI 04-FEB-2000
 DEFINITION hypothetical protein DKFZp434P1818.1 - human (fragment).
 ACCESSION T46372
 VERSION T46372 GI:11360168
 DBSOURCE pir: locus T46372;

 summary: #length 859 #checksum 5631
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 genetic: #note DKFZp434P1818.1
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 PIR dates: 04-Feb-2000 #sequence_revision 04-Feb-2000 #text_change
 04-Feb-2000
 ;
 punctuation in sequence.
 .
 KEYWORDS
 SOURCE Homo sapiens (human)
 ORGANISM Homo sapiens
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
 Hominidae; Homo.
 REFERENCE 1 (residues 1 to 859)
 AUTHORS Ottenwaelder,B., Obermaier,B., Mewes,H.W., Gassenhuber,J. and
 Wiemann,S.
 TITLE Direct Submission
 JOURNAL Submitted (??-JAN-2000) to the Protein Sequence Database
 COMMENT [WARNING] On Sep 13, 2005 this sequence was replaced by
 gi:74719265.
 FEATURES Location/Qualifiers
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 /db_xref="taxon:9606"
 Protein 1..859
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 Region <5..294
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 /note="Phosphoinositide polyphosphatase (Sac family)
 [Signal transduction mechanisms]; COG5329"
 /db_xref="CDD:34909"
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 181 frvncmdcld rtnvvqaaia rvvmegqlkk lgvmppeqpl pvkcnriyqi mwanngdsis
 241 rqyagtaalk gdftrtgerk lagvmkdgvn sanryylnrf kdayrqavid lmqqipvted
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361 thrdvdvlll lsnsayyvay yddevdkvnq yqrslenle kieigpeptl fgkpkfscmr
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481 kphediigir sqnqgslagg knflmskfss lnqkvqtkns nvnignlrkl gnftkpemkv
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781 hvtspsess ssravspfak irssmvqvas itqaglthgi nfavskvqks ppepeiinqv
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Sep 27 2006 15:22:06